IFW16

RAW SEQUENCE LISTING          DATE: 10/01/2004
PATENT APPLICATION:  US/09/920,137B     TIME: 10:46:27

Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt
Output Set: N:\CRF4\10012004\I920137B.raw

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       5 <110> APPLICANT: Giles-Komar, Jill
       6        David Shealy
       7        David Knight
       8        Bernie Scallon
       9        George Heavner
      11 <120> TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES
      13 <130> FILE REFERENCE: CEN0250
C-->  15 <140> CURRENT APPLICATION NUMBER: US/09/920,137B
C-->  15 <141> CURRENT FILING DATE: 2001-08-01
      15 <160> NUMBER OF SEQ ID NOS: 15
      17 <170> SOFTWARE: PatentIn Ver 3.1
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**ERRORED SEQUENCES**

Does Not Comply
Corrected Diskette Needed

(PS. 1-3)

← Pls see item #3 on error summary sheet.

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      19 <210> SEQ ID NO: 1
      20 <211> LENGTH: 5
      21 <212> TYPE: PRT
      22 <213> ORGANISM: Homo sapiens
W-->  23 <400> SEQUENCE: 1
      25  Arg Tyr Thr Met His
E-->  26  1   x       5   x
      28 <210> SEQ ID NO: 2
      29 <211> LENGTH: 17 -Found 16
      30 <212> TYPE: PRT
      31 <213> ORGANISM: Homo sapiens
W-->  32 <400> SEQUENCE: 2
      34  Val Ile Ser Phe Asp Gly Ser Asn Lys Tyr Tyr Val Asp Ser Val Lys
E-->  35  1   1           5   x               10  10          15  x
      37 <210> SEQ ID NO: 3
      38 <211> LENGTH: 10
      39 <212> TYPE: PRT
      40 <213> ORGANISM: Homo sapiens
W-->  41 <400> SEQUENCE: 3
      43  Glu Ala Arg Gly Ser Tyr Ala Phe Asp Ile
E-->  44  1   x           5   x           10  10
      64 <210> SEQ ID NO: 6
      65 <211> LENGTH: 10
      66 <212> TYPE: PRT
      67 <213> ORGANISM: Homo sapiens
W-->  68 <400> SEQUENCE: 6
      70  Gln Gln Arg Ser Asn Trp Pro Pro Phe Thr
E-->  71  1   x           5   5           10  x
```

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Input Set : **A:\Cen 0250 SEQ LIST 07-09-01.txt**
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      192 <210> SEQ ID NO: 13
      193 <211> LENGTH: (32)    Found 42
      194 <212> TYPE: DNA
      195 <213> ORGANISM: Homo sapiens
W--> 196 <400> SEQUENCE: 13
E--> 198   ctctcctgca gggccagtca gagtgttagc agctacttag cc
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                                                    32 — 42

**VERIFICATION SUMMARY**  DATE: 10/01/2004
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L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:23 M:283 W: Missing Blank Line separator, <400> field identifier
L:26 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:35 M:252 E: No. of Seq. differs, <211> LENGTH:Input:17 Found:16 SEQ:2
L:41 M:283 W: Missing Blank Line separator, <400> field identifier
L:44 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:50 M:283 W: Missing Blank Line separator, <400> field identifier
L:59 M:283 W: Missing Blank Line separator, <400> field identifier
L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:71 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:77 M:283 W: Missing Blank Line separator, <400> field identifier
L:108 M:283 W: Missing Blank Line separator, <400> field identifier
L:136 M:283 W: Missing Blank Line separator, <400> field identifier
L:172 M:283 W: Missing Blank Line separator, <400> field identifier
L:180 M:283 W: Missing Blank Line separator, <400> field identifier
L:188 M:283 W: Missing Blank Line separator, <400> field identifier
L:196 M:283 W: Missing Blank Line separator, <400> field identifier
L:198 M:254 E: No. of Bases conflict, LENGTH:Input:32 Counted:42 SEQ:13
L:198 M:252 E: No. of Seq. differs, <211> LENGTH:Input:32 Found:42 SEQ:13
L:204 M:283 W: Missing Blank Line separator, <400> field identifier
L:212 M:283 W: Missing Blank Line separator, <400> field identifier
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